



US009475853B2

(12) **United States Patent**  
**Hazlett et al.**

(10) **Patent No.:** **US 9,475,853 B2**  
(45) **Date of Patent:** **Oct. 25, 2016**

(54) **FUSION PROTEIN FOR ENHANCING  
IMMUNOGENICITY OF BACTERIAL  
ANTIGEN/IMMUNOGEN**

(58) **Field of Classification Search**

None

See application file for complete search history.

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(\*) Notice: Subject to any disclaimer, the term of this  
patent is extended or adjusted under 35  
U.S.C. 154(b) by 0 days.

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(21) Appl. No.: **14/373,212**

(22) PCT Filed: **Jan. 22, 2013**

(86) PCT No.: **PCT/US2013/022526**

§ 371 (c)(1),

(2) Date: **Jul. 18, 2014**

*Primary Examiner* — Brian J Gangle

(87) PCT Pub. No.: **WO2013/110064**

PCT Pub. Date: **Jul. 25, 2013**

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(65) **Prior Publication Data**

US 2015/0030632 A1 Jan. 29, 2015

(57) **ABSTRACT**

Establishment of an effective and uniform vaccine develop-  
ment strategy is key to conquering current and emerging  
infectious diseases. Despite successes against an array of  
bacterial agents, current approaches to vaccine development  
are as diverse as the microbes they target and require  
adjuvants that often have limited efficacy and/or toxic side  
effects. As a consequence, vaccine discovery is often slow,  
inefficient, and unsuccessful in the case of many high  
priority pathogens. The present disclosure suggests that  
vaccine generation for bacterial pathogens can be improved  
by optimizing the efficiency of processing/presentation of a  
bacterial immunogen via the targeting of immunogen to  
CR2 and/or TLR2 on APCs. This approach not only yields  
an adjuvant-free mucosal vaccine against a Category A  
biothreat agent, but also establishes a novel genetic  
approach/platform for vaccine development, which is appli-  
cable to many other infectious agents, thereby profoundly  
impacting preventive medicine/public health.

#### Related U.S. Application Data

(60) Provisional application No. 61/588,368, filed on Jan.  
19, 2012, provisional application No. 61/681,996,  
filed on Aug. 10, 2012.

(51) **Int. Cl.**

**A61K 39/02** (2006.01)

**C07K 14/47** (2006.01)

**C07K 14/195** (2006.01)

**A61K 39/00** (2006.01)

(52) **U.S. Cl.**

CPC ..... **C07K 14/472** (2013.01); **A61K 39/0208**  
(2013.01); **A61K 39/0291** (2013.01); **C07K**  
**14/195** (2013.01); **A61K 2039/53** (2013.01);  
**A61K 2039/6031** (2013.01); **C07K 2319/00**  
(2013.01); **C07K 2319/33** (2013.01)

**27 Claims, 6 Drawing Sheets**

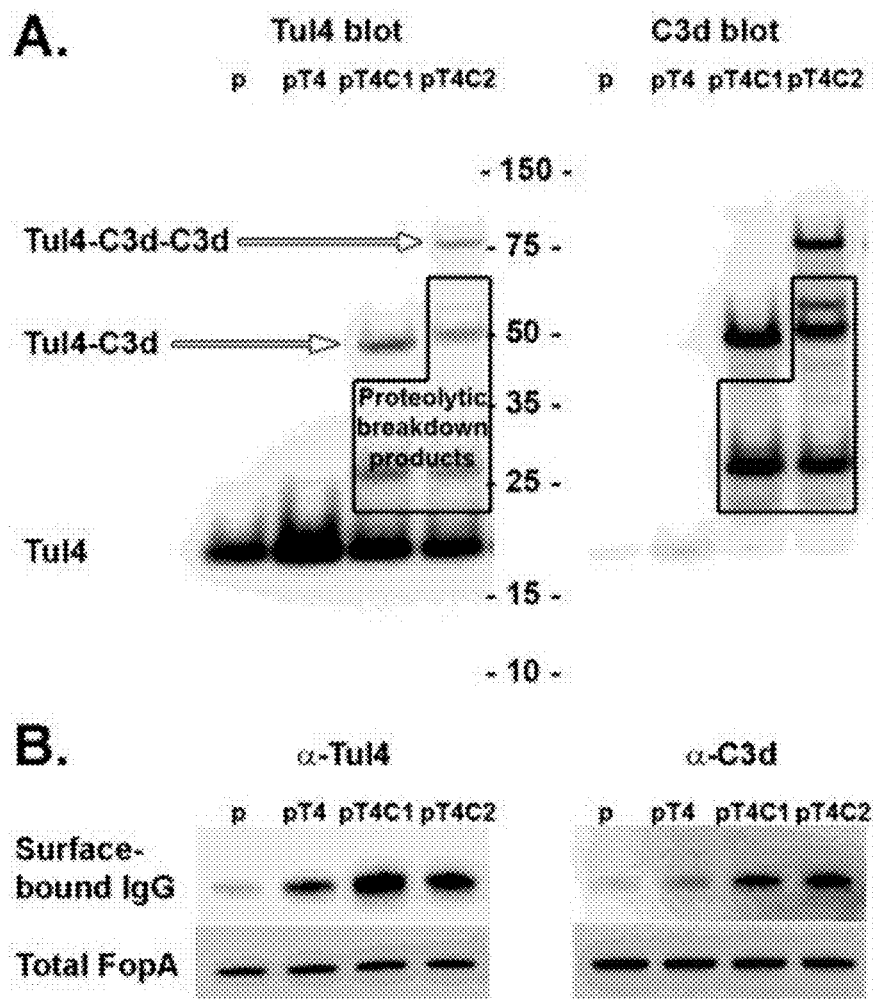


FIGURE 1

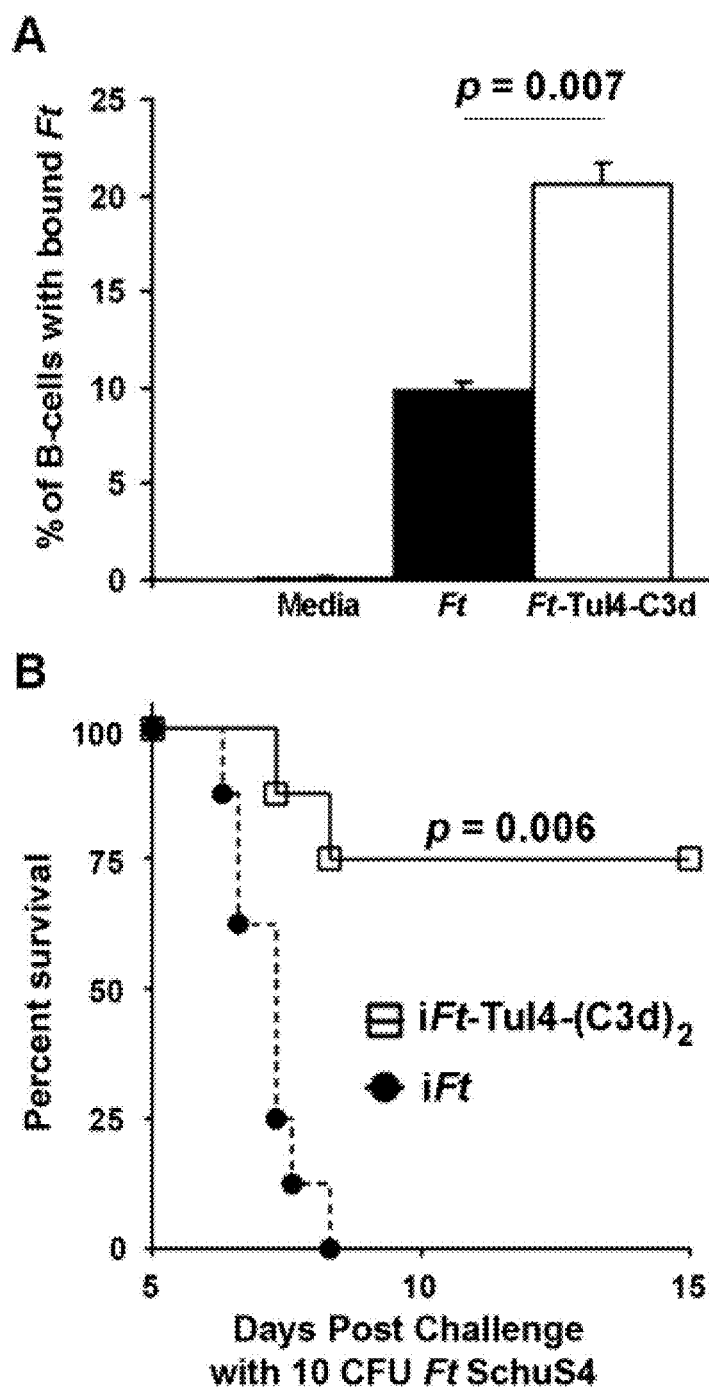


FIGURE 2

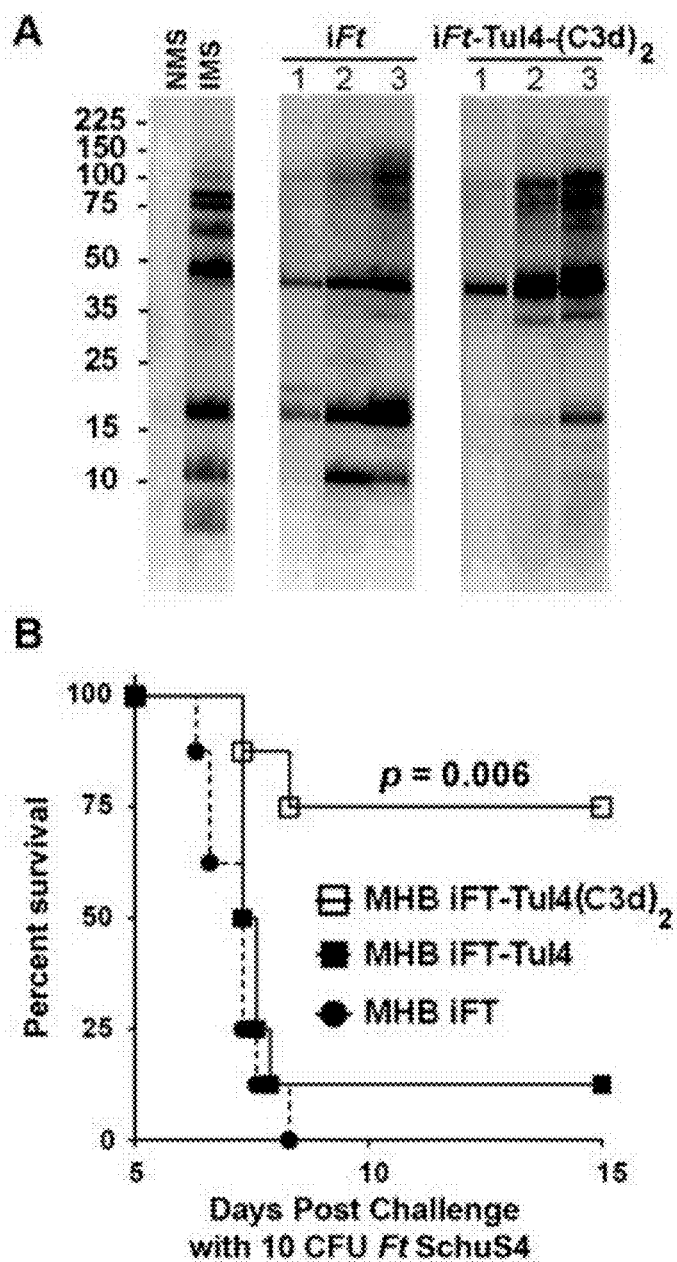
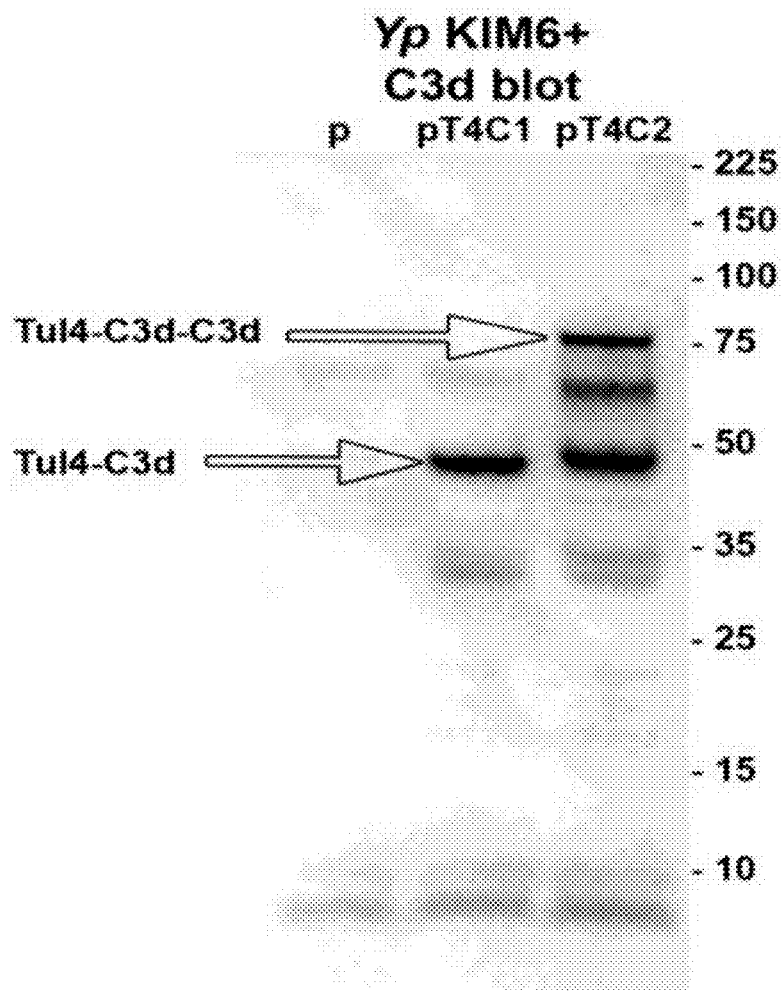


FIGURE 3

**FIGURE 4**

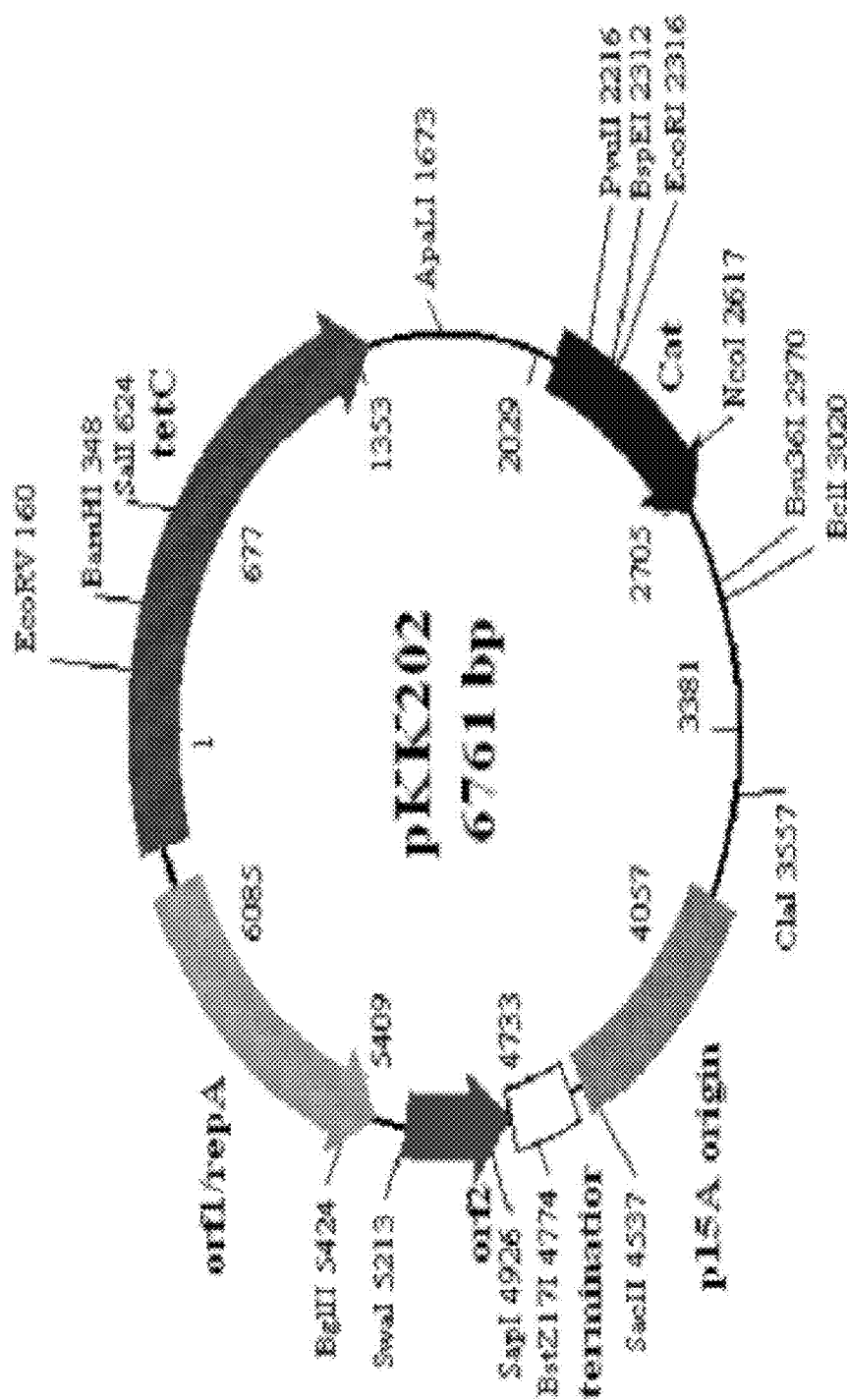


FIGURE 5

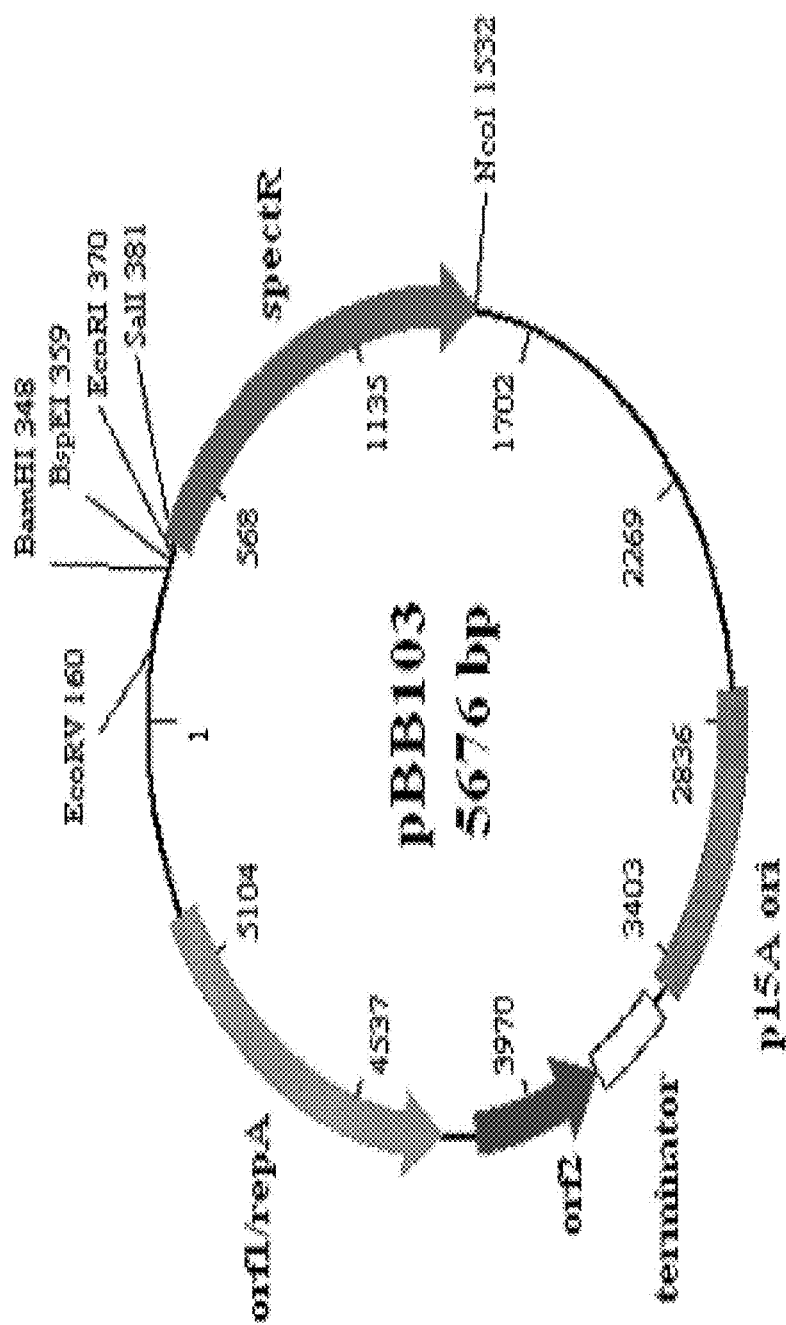


FIGURE 6

# FUSION PROTEIN FOR ENHANCING IMMUNOGENICITY OF BACTERIAL ANTIGEN/IMMUNOGEN

## CROSS REFERENCE TO RELATED APPLICATIONS

This application is a 371 National Phase of PCT International Application No. PCT/US2013/022526 filed on Jan. 22, 2013, and published in English as WO 2013/110064 A1 on Jul. 25, 2013, which claims priority to U.S. provisional application Ser. Nos. 61/588,368, filed Jan. 19, 2012 and 61/681,996, filed Aug. 10, 2012. The contents of these disclosures are hereby incorporated by reference into the present application.

## GOVERNMENT RIGHTS STATEMENT

This invention was made with U.S. Government support under grant no. PO1-AI056320 awarded by the NIAID of the National Institutes of Health. The U.S. Government has certain rights in the invention.

## SEQUENCE LISTING

This application contains a Sequence Listing, created on Jan. 15, 2013; the file, in ASCII format, is designated 0410046AWO\_sequencelisting\_ST25.txt and is 13.8 kilobytes in size. The sequence listing file is hereby incorporated by reference in its entirety into the application.

## TECHNICAL FIELD

The present disclosure relates generally to vaccine development. More particularly, the present disclosure relates to a polynucleotide construct that enhances immunogenicity of a bacterial pathogen transformed with the construct.

## BACKGROUND OF THE DISCLOSURE

*Francisella tularensis* (Ft)) is a Gram-negative, facultative intra-cellular pathogen and the etiological agent of tularemia, a disease that can be fatal in humans. Ft can infect individuals at both mucosal and peripheral sites, with the former being the most lethal form of infection. Both humoral and cellular immunity play a role in protection against this organism. While there are only about 200 known Ft infections per year in the U.S., as few as one organism can be fatal, if inhaled. This has led to the obvious concern that Ft could be used as a bioterrorism agent, when dispersed as an aerosol in a populated area. Thus, Ft has been listed as a category A select agent, and there is strong interest in the development of an effective vaccine against Ft.

Furthermore, the development of needle-free, mucosal vaccines that do not require exogenous adjuvants is of significant interest to all areas of infectious disease including biodefense. Even though some infections can be treated successfully with antibiotics, under some circumstances, such as biodefense, vaccination is preferred over post-exposure therapy.

To date, the current gold standard for tularemia prophylaxis, infection with viable Live Vaccine Strain (LVS) Ft, provides only moderate protection in humans. Thus, a need exists for a vaccine that yields improved efficacy and a major effort is underway to develop a new generation of safe and effective vaccines against the disease.

## SUMMARY OF THE DISCLOSURE

The present disclosure is directed to fusion proteins and nucleic acids encoding those fusion proteins, which, when expressed in a pathogenic bacterial cell that is administered to a subject, significantly enhances immune responsiveness in the subject to the pathogen.

In one aspect, therefore, the disclosure relates to fusion proteins and nucleic acids that encode the fusion proteins, the fusion protein comprising a *Francisella tularensis* (Ft) lipoprotein such as Tul4A that binds toll-like receptor 2 (TLR2) linked to a portion of complement component 3 (C3), such as C3d, that binds to a complement receptor type 2 (CR2) on immune effector cells. The nucleic acid(s) may be part of a polynucleotide that comprises an expression cassette, that is, it may be part of a construct that includes a promoter and/or other regulatory elements that effect expression. Vectors which include the nucleic acid and/or an expression cassette containing the nucleic acid are encompassed by the disclosure.

In another aspect, therefore, the disclosure relates to a polynucleotide, for example, a vector comprising a nucleic acid that encodes a fusion protein comprising a Ft lipoprotein that binds toll-like receptor 2 (TLR2) such as Tul4A linked to C3d. The nucleic acid encoding the fusion protein or expression cassette incorporating the nucleic acid encoding the fusion protein can be cloned into a variety of plasmids to achieve expression of the fusion protein in a wide range of bacteria.

In yet another aspect, the disclosure relates to a bacterial cell that has been transformed using a vector comprising a nucleic acid encoding a fusion protein of a Ft lipoprotein that binds toll-like receptor 2 (TLR2) such as Tul4A linked to C3d. The transformed bacterial cell expressing the fusion p

In a related aspect, the disclosure relates to a pharmaceutical composition comprising a bacterial cell that has been transformed with a vector comprising a nucleic acid encoding a fusion protein of Tul4A linked to C3d. The transformed bacterial cell expresses the fusion protein, which is anchored to the outer membrane of the bacteria.

## BRIEF DESCRIPTION OF THE DRAWINGS

FIG. 1A-B shows expression and surface-exposure of Tul4A and C3d by engineered Ft LVS. A) Ft strains harboring either the empty vector (p), the Tul4A over-expression vector (pT4), or the Tul4A-C3d vectors (pT4C1 and pT4C2) were probed by western blot for total Tul4A and C3d. In B, intact bacteria were incubated with  $\alpha$ -Tul4A or  $\alpha$ -C3d Ab, washed, and probed by western blot for IgG heavy chain as an indication of surface-exposed Tul4A or C3d. The blot was subsequently stripped and re-probed for total FopA as a loading control.

FIG. 2 shows that engineered expression of Tul4A-C3d by Ft promotes binding to CR2-expressing cells. CR2-positive B cells (A20 cells) were incubated for 2 hrs with media or fluorescently-labeled (pKH26-treated) Ft LVS strains. Following fixation and washing, the B cells were analyzed by flow cytometry.

FIG. 3A-B shows that the immunogenicity and protective capacity of iFt LVS immunogens is enhanced by endogenous Tul4A-C3d production. C57B/6 mice (8 per group) were immunized and boosted i.n. with  $4 \times 10^7$  iFt LVS (WT, Tul4A, or Tul4A-C3d-C3d) per mouse on days 0, 14, and 28. A) Pooled test sera, drawn 7 days after each immunization (#1, 2, and 3), was used in western blot analysis at a 1:500 dilution to probe BHI-grown Ft LVS; normal mouse sera



(NMS) and infection-derived immune mouse sera (IMS) were used as controls. B) On day 42, mice were challenged i.n. with Ft SchuS4. Survival was monitored three times daily for 21 days. No deaths occurred after day 8; PBS-immunized control mice succumb by day 7.

FIG. 4A-B shows expression of Tul4A-C3d(n) produced by recombinant *Yersinia pestis* (Yp). Bacterial lysates from Yp KIM6<sup>+</sup> strains harboring either the empty vector (p), or the Tul4A-C3d vectors (pT4C1+pT4C2) were resolved by SDS-PAGE and probed by western blot with antibodies which recognize the 17 kDa Ft lipoprotein Tul4AA or the 33 kDa murine complement component 3 fragment, C3d. Plasmids are designated as follows: "p" is empty vector (pBB103); "pT4" is pBB103 expressing Tul4AA; "pT4Cx" is "pT4" with one ("pT4C1") or two ("pT4C2") copies of C3d linked sequentially to the C-terminus of Tul4AA.

FIG. 5 is a schematic of plasmid pKK202.

FIG. 6 is a schematic of plasmid pBB103.

#### DETAILED DESCRIPTION OF THE DISCLOSURE

All patents, published applications and other references cited herein are hereby incorporated by reference in their entirety.

In practicing the present disclosure, many conventional techniques in microbiology, molecular biology and immunology are used, which are within the skill of the ordinary artisan. Some techniques are described in greater detail in, for example, *Molecular Cloning: a Laboratory Manual* 3rd edition, J. F. Sambrook and D. W. Russell, ed. Cold Spring Harbor Laboratory Press 2001, the contents of this and other references containing standard protocols, widely known to and relied upon by those of skill in the art, including manufacturers' instructions are hereby incorporated by reference as part of the present disclosure.

Abbreviations used herein include the following:

APCs: antigen presenting cells

BMDs: bone marrow dendritic cells

BMDM: bone-marrow-derived macrophages

C3d: complement component 3 fragment d

FDCs: follicular dendritic cells

Ft: *Francisella tularensis*

Ft LVS: *F. tularensis* live vaccine strain

TLR2: toll-like receptor 2

Tul4A: a *F. tularensis* membrane lipo-protein; also known as LpnA or FTT0901.

Yp: *Yersinia pestis*

Unless otherwise defined, all technical and scientific terms used herein have the same meaning as commonly understood by one of ordinary skill in the art to which this disclosure belongs. The terminology used in the description of the disclosure herein is for the purpose of describing particular embodiments only and is not intended to be limiting of the disclosure. All publications, patent applications, patents, and other references mentioned herein are incorporated by reference in their entirety.

The term "fusion polypeptide" or "fusion protein" refers to a protein having at least two heterologous polypeptides covalently linked, either directly or via an amino acid linker. The polypeptides forming the fusion protein are typically linked C-terminus to N-terminus, although they can also be linked C-terminus to C-terminus, N-terminus to N-terminus, or N-terminus to C-terminus. The polypeptides of the fusion protein can be in any order and may include more than one of either or both of the constituent polypeptides. The term encompasses conservatively modified variants, polymorphic

variants, alleles, mutants, subsequences, interspecies homologs, and immunogenic fragments of the antigens that make up the fusion protein. Fusion proteins of the disclosure can also comprise additional copies of a component antigen or immunogenic fragment thereof.

A "recombinant expression cassette" or simply an "expression cassette" is a nucleic acid construct, generated recombinantly or synthetically, with nucleic acid elements that are capable of affecting expression of a structural gene in hosts compatible with such sequences. Expression cassettes include at least promoters and optionally, transcription termination signals. Typically, the recombinant expression cassette includes a nucleic acid to be transcribed (e.g., a nucleic acid encoding a desired polypeptide), and a promoter. Additional factors necessary or helpful in effecting expression may also be used as described herein. An expression cassette can be designed to include or delete transcription termination signals, enhancers, and other nucleic acid sequences that influence gene expression to achieve a desired expression product. In some embodiments, a recombinant expression cassette encoding an amino acid sequence for a fusion protein of the disclosure is expressed in a bacterial host cell.

The term "operably linked" refers to functional linkage between a nucleic acid expression control sequence (such as a promoter, signal sequence, or array of transcription factor binding sites) and a second nucleic acid sequence, wherein the expression control sequence affects transcription and/or translation of the nucleic acid corresponding to the second sequence.

The term "heterologous" when used with reference to portions of a nucleic acid indicates that the nucleic acid comprises two or more subsequences that are not found in the same relationship to each other in nature. For instance, the nucleic acid is typically recombinantly produced, having two or more sequences from unrelated genes arranged to make a new functional nucleic acid, e.g., a promoter from one source and a coding region from another source, or coding regions from different sources. Similarly, a heterologous protein indicates that the protein comprises two or more subsequences that are not found in the same relationship to each other in nature (e.g., a fusion protein).

The term "linker" or "linked" refers to the covalent linkage between two polypeptides in a fusion protein. The polypeptides are typically joined via a peptide bond, either directly to each other or via one or more additional amino acids. The composition and length of the linker may be determined in accordance with methods well known in the art and may be tested for efficacy. The linker is generally from about 3 to about 15 amino acids long, in some embodiments about 5 to about 10 amino acids long, however, longer or shorter linkers may be used or the linker may be dispensed with entirely. A glycine linker is one that contains one or more glycines but no other amino acids, e.g. GGGG (SEQ ID NO: 14). A glycine-rich linker is one that contains one or more glycines and may contain other amino acids as long as glycine is the predominant species in the linker e.g. GGGNGG (SEQ ID NO: 13). A glycine-serine linker is one which contains both glycine and serine in any proportion, e.g. GGGS (SEQ ID NO: 11).

The term "adjuvant" refers to the components in a vaccine or therapeutic composition that increase the specific immune response to the antigen (see, e.g., Edelman, AIDS Res. Hum Retroviruses 8:1409-1411 (1992)). Adjuvants are well known to those of skill in the art and may include cytokines (e.g., IFN- $\gamma$ , IL-2, and IL-12) which contribute to the induc-

tion of cell-mediated immune response to an administered antigen, as well as induction of humoral immune responses.

The terms “nucleic acid” and “polynucleotide” refer to deoxyribonucleotides and polymers thereof.

Unless otherwise indicated, a particular nucleic acid sequence also implicitly encompasses conservatively modified variants thereof (e.g., degenerate codon substitutions) and complementary sequences, as well as the sequence explicitly indicated. Specifically, degenerate codon substitutions may be achieved by generating sequences in which the third position of one or more selected (or all) codons is substituted with mixed-base and/or deoxyinosine residues (Batzer et al., *Nucleic Acid Res.* 19:5081 (1991); Ohtsuka et al., *J. Biol. Chem.* 260:2605-2608 (1985); Rossolini et al., *Mol. Cell. Probes* 8:91-98 (1994)). The term nucleic acid is used interchangeably with cDNA, mRNA, oligonucleotide, and polynucleotide.

Furthermore, one of skill will recognize that individual substitutions, deletions or additions which alter, add or delete a single amino acid or a small percentage of amino acids (typically less than 5%, more typically less than 1%) in an encoded sequence are “conservatively modified variations” where the alterations result in the substitution of an amino acid with a chemically similar amino acid. Conservative substitution tables providing functionally similar amino acids are well known in the art.

The terms “polypeptide,” and “protein” are used interchangeably herein to refer to a polymer of amino acid residues. The terms encompass amino acid polymers in which one or more amino acid residue is an artificial chemical mimetic of a corresponding naturally occurring amino acid, as well as to naturally occurring amino acid polymers and non-naturally occurring amino acid polymer.

The term “amino acid” refers to naturally occurring and synthetic amino acids, as well as amino acid analogs and amino acid mimetics that function in a manner similar to the naturally occurring amino acids. Naturally occurring amino acids are those encoded by the genetic code, as well as those amino acids that are later modified, e.g., hydroxyproline, -gamma-carboxyglutamate, and O-phosphoserine. Amino acid analogs refers to compounds that have the same basic chemical structure as a naturally occurring amino acid, i.e., an alpha carbon that is bound to a hydrogen, a carboxyl group, an amino group, and an R group, e.g., homoserine, norleucine, methionine sulfoxide, methionine methyl sulfonium. Such analogs have modified R groups (e.g., norleucine) or modified peptide backbones, but retain the same basic chemical structure as a naturally occurring amino acid. Amino acid mimetics refers to chemical compounds that have a structure that is different from the general chemical structure of an amino acid, but that functions in a manner similar to a naturally occurring amino acid.

Amino acids may be referred to herein by either their commonly known three letter symbols or by the one-letter symbols recommended by the IUPAC-IUB Biochemical Nomenclature Commission. Nucleotides, likewise, may be referred to by their commonly accepted single-letter codes.

As to amino acid sequences, one of skill will recognize that individual substitutions, deletions or additions to a nucleic acid, peptide, polypeptide, or protein sequence which alters, adds or deletes a single amino acid or a small percentage of amino acids in the encoded sequence is a “conservatively modified variant” where the alteration results in the substitution of an amino acid with a chemically similar amino acid. Conservative substitution tables providing functionally similar amino acids are well known in the art. Such conservatively modified variants are in addition to

and do not exclude polymorphic variants, interspecies homologs, and alleles of the disclosure.

The present disclosure is based, in part, on the observation that the expression by a bacterium of a fusion protein combining a TLR2 ligand (e.g. Tul4A) and a complement receptor ligand (e.g. C3d), can (1) increase binding of the bacterium by cells bearing CR2 such as B-cells or FDCs; and (2) increase survival of animals that were immunized with bacteria bearing the fusion protein on their surface prior to challenge with the bacteria. The fusion protein disclosed herein, therefore, enhances immunogenicity of a whole bacteria immunogen when the fusion protein is expressed at the surface, thereby boosting the immune response mounted in a subject against that bacteria.

Without wishing to be bound by theory, the expression of a Tul4A-C3d(n) fusion protein of the disclosure by a bacterial cell enhances the immunogenicity of that bacterial immunogen by targeting of the bacteria (inactivated for example) to host cells bearing the complement receptor CR2 while at the same time stimulating TLR2 on the same host immune effector cells at an increased level due to the overexpression of Tul4A. This results in highly-efficient delivery of the immunogen (the bacteria expressing Tul4A-C3d(n)) to the most potent antigen-presenting cells that, by virtue of TLR2 stimulation, should become highly activated to initiate a robust immune response.

Tul4A is an integral membrane lipoprotein expressed across different strains of *Francisella*. In engineering bacterial cells that express a Tul4A-C3d(n) fusion protein, transformation with a nucleic acid of the invention encodes a fusion protein that is not secreted but membrane-bound.

Tul4A binds to the host toll-like receptor 2 (TLR2) to activate cells bearing this receptor. Expression of a Tul4A-C3d(n) fusion protein increases the concentration of Tul4A on the surface of the bacterial cell; Tul4A acts as a protein adjuvant for antigen (Ag)—presenting cells (APCs). Additionally, TLR2 ligands may also target delivery of antigens to these APCs.

C3d is a fragment of mammalian complement component 3 which binds the complement receptor 2 (CR2) expressed on B-cells and follicular dendritic cells (FDCs); i.e. C3d targets CR2 for delivery of C3d-linked moieties to B-cells and FDCs. Linking Tul4A-C3d fusion proteins to whole cell immunogens targets delivery of the immunogen to CR2-positive cells (and potentially TLR2-positive cells) and stimulate a potent anti-immunogen response through TLR2-stimulation. Since the immunogen is targeted (via C3d) and adjuvanted (via Tul4A), the need for external adjuvants should be reduced/eliminated.

Plasmids

In one embodiment, coding regions of the *Francisella tularensis* (Ft) gene encoding Tul4A (a surface-exposed lipoprotein) (Genbank accession no. FTT0901) were genetically fused in-frame with DNA encoding residues 1024-1320 from murine C3. C3 encoding DNA was PCR amplified from plasmid pMLC3/7 (ATCC, Manassas, Va.) and fused in-frame to the 3' end of either Tul4A or PilA (the SchuS4-specific pilin subunit protein.) This portion of murine C3 is 91% identical to human C3d. Both of these Ft genes lack a stop codon to allow for the production of Tul4A-C3d and/or PilA-C3d fusion proteins.

In some embodiments, constructs which encode multiple copies of C3d (these are designated C3d(n) where n=an integer from 1 to 5) downstream of Tul4A are desirable; these were developed to increase the avidity of the Tul4A-C3d(n) chimeras for CR2.

In one embodiment, the chimeric gene is placed downstream of a promoter, such as a groEL promoter and the groEL:Tul4A-C3d(n) gene fusion is moved into a plasmid(s) that will replicate and facilitate expression of the fusion protein. Useful expression vectors for bacterial hosts include bacterial plasmids, such as pBB103 and pMP633 and wider host range plasmids, such as pBAV1k-T5-gfp (Addgene, Cambridge Mass.), and pSE100 (Addgene, Cambridge Mass.). Nucleic acids encoding the fusion protein or expression cassettes incorporating the nucleic acid encoding the fusion protein can therefore, be cloned into a variety of plasmids to achieve expression of the fusion protein in a wide range of bacteria including *Francisella tularensis*, *Escherichia coli*, *Yersinia pestis*, *Acinetobacter baumannii*, *Mycobacterium tuberculosis*, *Borrelia burgdorferi*, *Methicillin-resistant Staphylococcus aureus*, *Klebsiella pneumonia*, Groups A and B. *streptococcus*, *Pseudomonas aeruginosa*, *Vancomycin-Resistant Enterococci*, *Multidrug-Resistant Neisseria gonorrhoeae*, *Helicobacter pylori*, and *Bartonella henselae*.

In one embodiment, a plasmid, designated pBB103, was created by inserting a small MCS and the spectinomycin resistance gene into BamHI/NcoI of pKK202. This effectively replaced most of the tet and chlor resistance genes in pKK202, however, the tetC promoter remains and drives expression of whatever is cloned into the pBB103 MCS (provided it is in the same orientation as PtetC). A schematic of each of pKK202 and pBB103 is shown in FIGS. 5 and 6, respectively, and the nucleotide sequence of pBB103 is given in SEQ ID NO: 4.

*F. tularensis* (Ft) and *Y. pestis* (Yp) were transformed with a plasmid containing the expression cassette incorporating a nucleic acid sequence that encodes the fusion protein using methods known to those of skill in the art. Following transformation, Ft and Yp strains harboring either the empty vector (p), the Tul4A over-expression vector (pT4), or the Tul4A-C3d and Tul4A-C3d-C3d vectors (pT4C1 and pT4C2, respectively) were probed by western blot for total Tul4A and C3d. Intact bacteria were incubated with  $\alpha$ -Tul4A or  $\alpha$ -C3d antibody (Ab), washed, and probed by western blot for IgG heavy chain as an indication of surface-exposed Tul4A or C3d. The blot was subsequently stripped and re-probed for total FopA as a loading control. FIGS. 1A and 1B shows expression and surface-exposure of Tul4A and C3d by engineered Ft LVS.

Yp KIM6<sup>+</sup> were similarly transformed and evaluated for expression of the fusion protein. Briefly, bacterial lysates from Yp KIM6<sup>+</sup> strains harboring either the empty vector (p), or the Tul4A-C3d vectors (pT4C1+pT4C2) were electrophoresed and probed by western blot with antibodies which recognize the 17 kDa Ft lipoprotein Tul4AA or the 33 kDa murine complement component 3 fragment, C3d. Plasmids are designated as follows: "p" is empty vector (pBB103); "pT4" is pBB103 expressing Tul4AA; "pT4Cx" is "pT4" with one ("pT4C1") or two ("pT4C2") copies of C3d linked sequentially to the C-terminus of Tul4AA. As shown in FIG. 4, transformation of these cells resulted in expression of Tul4A-C3d(n) by recombinant Yp.

Next, to determine whether bacteria bearing the Tul4A-C3d/Tul4A-C3d-C3d fusion protein would be bound by B cells, CR2-positive B cells (A20 cells) were incubated for 2 hrs with media or fluorescently-labeled (pKH26-treated) Ft LVS strains. Following fixation and washing, the B cells were analyzed by flow cytometry. FIG. 2 shows that engineered expression of Tul4A-C3d by Ft promotes binding to CR2-expressing cells.

To determine whether immunogenicity and protective capacity of iFt LVS immunogens was enhanced by fusion protein expression, C57B/6 mice (8 per group) were immunized and boosted intranasally (i.n.) with  $4 \times 10^7$  iFt LVS (wild type (WT), Tul4A, or Tul4A-C3d-C3d) per mouse on days 0, 14, and 28. Pooled test sera, drawn 12 days after each immunization (#1, 2, and 3), was used in western blot analysis at a 1:500 dilution to probe BHI-grown Ft LVS; normal mouse sera (NMS) and infection-derived immune mouse sera (IMS) were used as controls. On day 42, mice were challenged i.n. with Ft SchuS4. Survival was monitored three times daily for 21 days.

As shown in the Kaplan-Meier plot in FIG. 3B, no deaths of animals immunized using the fusion protein occurred after day 8; PBS-immunized control mice succumb by day 7. Furthermore, even though overexpression of Tul4A alone conferred some benefit, the survival of animals immunized with iFt Tul4A-C3d-C3d was significantly improved, showing that the immunogenicity and protective capacity of iFt LVS immunogens is enhanced by endogenous Tul4A-C3d production.

#### Advantages of the Technology

The development of needle-free, mucosal vaccines that do not require exogenous adjuvants is of significant interest to all areas of infectious disease including biodefense. By combining CR2-targeting with TLR2-stimulation the presently disclosed approach addresses this need. Furthermore, as this approach genetically/physically-links the targeting and stimulation moieties to the immunogen, no further manipulation (addition of exogenous targeting moieties, external adjuvants, etc.) of the prepared immunogen is required.

## EXAMPLES

### Construction of Genetic—Targeting Vectors, pT4C(n)

Using techniques known to those of skill in the art, DNA encoding the surface-exposed lipoprotein (2) Tul4AA (FTT0901) from *Francisella tularensis* (Ft) SchuS4 was PCR-amplified from the bacterium's genome using primers:

5' EcoR1SDTul4A: (SEQ ID NO: 5)  
GAA TTC TGA ATA TTT AAA AAT AGG AGT ATC TAT ATG  
and

3' BamH1-Tul4A: (SEQ ID NO: 6)  
GGA TCC TCA TTA AAT ATT TAT TGA ATC AGA AGC GAT

TAC

and subsequently cloned into TOPO PCR 2.1 (Invitrogen). Following validation, the Tul4A insert was sub-cloned as an EcoR1/BamH1 fragment into the same sites of plasmid pF (1) which contains the strong Ft groEL promoter (PgroEL) driving expression of the insert, in this case Tul4A. The PgroEL-Tul4A fusion was amplified from the above plasmid using:

forward primer 5' PgroE-KpnBg12: (SEQ ID NO: 7)  
GGT ACC AGA TCT TTG TAT GGA TTA GTC GAG CTA AAA  
AGC

-continued

and either of the

reverse primers 3' BamHI-Tul4A or  
3' BamHI-Tul4A-NS:

(SEQ ID NO: 8) 5  
GGA TCC CCC TCC AAT ATT TAT TGA ATC AGA AGC GAT

TAC

and cloned into TOPO PCR 2.1. The 3' BamHI-Tul4A-NS primer eliminates the native Tul4A stop codon allowing for construction of translational fusions with this lipoprotein which is a known TLR2 agonist (3). These two PgroEL-Tul4A fusions (with and w/o Tul4A stop codons) were digested out of the TOPO PCR 2.1 backbone as Bgl2/XhoI fragments and ligated into the BamHI/SalI sites of pBB103 resulting in plasmids pT4 which contains the Tul4A stop codon, and pT4NS which lacks the stop codon and instead contains a GGGs "1/2 linker" at the C-terminus of Tul4A.

DNA encoding residues 1011 to 1293 (prepro-C3 numbering) of murine C3d was amplified from plasmid pMLC3/7 (ATCC, Manassas, Va.) using the following primers:

5' C3d Bgl2: (SEQ ID NO: 9) 25  
AGA TCT GGG GGA GGC TCT GGG GAA CAG AAC ATG  
and

3' C3d BSSB: (SEQ ID NO: 10)  
AGA TCT GTC GAC TCA GGA TCC ACC TCC GTT CAA GTC

CTT ATG GTC AGG GAC;

the product was cloned into TOPO PCR 2.1. This 5' primer encodes a GGGs (SEQ ID NO: 11) "1/2 linker" at the N-terminus immediately upstream of Gly 1011 of C3d. The 3' primer encodes a GGGs (SEQ ID NO: 11) "1/2 linker" at the C-terminus of C3d immediately downstream of Asn 1293; the primer also contains a BamHI site followed by a stop codon followed by a SalI site followed by a Bgl2 site. Following validation, the c3d insert was sub-cloned as a Bgl2 fragment into the BamHI site of plasmid pT4NS and a clone in which the c3d insert was similarly oriented to the up-stream Tul4A gene was selected. This plasmid, in which the groEL promoter drives expression of the Tul4A-GGGSGGGs (SEQ ID NO: 12)-C3d-GGGs (SEQ ID NO: 11) fusion protein, was dubbed pT4C1. To construct pT4C2, which contains two tandemly arranged copies of the c3d insert, pT4C1 was digested with BamHI and SalI and ligated to fresh c3d insert as a Bgl2/SalI fragment. The same process, albeit starting with pT4C2 instead of pT4C1, was used to generate pT4C3.

Bacteria and Media

Bacteria were grown and maintained in accordance with methods known in the art. Wild-type (WT) *F. tularensis* LVS has been described previously [12], [36], [114]. *F. tularensis* SchuS4, originally isolated from a human case of tularemia, was obtained from the U.S. Army Medical Research Institute for Infectious Diseases (Frederick, Md.). All experiments using SchuS4 were conducted within the Albany Medical College ABSL-3/BSL-3 facility which has been certified by the Center for Disease Control. SchuS4 lysates for protein analysis to be performed outside of this facility were generated by boiling harvested bacteria in 50 mM Tris, pH 8.0 containing 1% SDS followed by sterility testing.

Routine culturing of *F. tularensis* involved streaking aliquots of frozen bacterial glycerol stocks onto Mueller Hinton Chocolate agar plates (Becton, Dickinson and Com-

pany (BD), Sparks, Md.) followed by 2-3 days of growth in a humidified chamber maintained at 37° C. Starter cultures were generated by resuspending several isolated colonies in ~100 µl of BHI from which ~50 µl was immediately used to inoculate 3-5 ml of BHI and MHB for ~18 hrs of growth on an orbital shaker operating at 200 rpm and 37° C. Mature starter cultures were used at a 1:100 dilution to inoculate larger volumes (10-50 ml within 125-250 ml Erlenmeyer flasks) of fresh media appropriate for each experiment. Unless stated otherwise, all bacterial cultures used in this work were in mid-log phase (OD<sub>600</sub>=0.4-0.6). Preparation of MHB and BHI media as well as supplementation of BHI with casamino acids has been described previously [12]. Supplementation of MHB with spermine to 200 µM was performed as described [34].

SDS-PAGE and Western Blot Analysis

Routine SDS-PAGE characterization was used to assess the integrity of the fusion proteins. Samples of *Francisella* [10 µg (~1×10<sup>8</sup> cells)] were mixed with Laemmli sample buffer and boiled for 10 min prior to resolution through 4-12% gradient SDS-PAGE pre-cast gels (Invitrogen). The running buffer was NuPAGE MES SDS buffer from Invitrogen; gels were variously run at 90-160 V. Resolved gels were stained with either coomassie blue (BioRad) or transferred to nitrocellulose membranes. Coomassie-stained gels were scanned into Adobe Photoshop using an HP 2820. Membranes to be probed with mAb were blocked for 15 min with PBS, 0.05% Tween Polysorbate 20 (TWEEN 20), 1% casein; blots to be probed with polyclonal sera were blocked for 1 hr with PBS, 0.05% TWEEN 20, 2.5% horse serum, 1% casein. Polyclonal sera were applied for 1-2 hr at dilutions ranging from 1:1000 to 1:60,000. Supernatants from mAb-producing hybridomas were applied for overnight (o/n) incubations. The mAb FB11 specific for *F. tularensis* O-antigen (Abcam) was used o/n at a dilution of 1:1000. Blots to be probed multiple times were first probed with mAb prior to stripping and reprobing with polyclonal sera. HRP-conjugated secondary antibodies were used at dilutions ranging from 1:1000 to 1:20,000. Development of the chemiluminescent substrate (SuperSignal West Pico, Pierce, Rockford, Ill.) was visualized using an Alpha Innotech imaging system in movie mode. Densitometric analysis of developed blots was performed on the same system.

Staining of SDS-PAGE Gels

Boiled samples of *F. tularensis* in Laemmli sample buffer (1 µg/ul protein) were de-proteinized with proteinase K (0.4 µg/ul final) at 60° C. for 1 hour and re-boiled prior to resolution through 4-12% gradient SDS-PAGE pre-cast gels (Invitrogen). Each lane was loaded with the de-proteinized material generated from *F. tularensis* lysate containing 10 µg of protein. Carbohydrates were visualized in situ using the Pro-Q Emerald 300 Lipopolysaccharide Gel Stain Kit (Invitrogen) as instructed. Briefly, resolved gels were fixed overnight in 5% acetic acid, 50% methanol. Following two 20 minute washes in 3% acetic acid, the gels were incubated for 30 minutes in oxidizing solution, washed 3 times in 3% acetic acid prior to fluorescent staining of oxidized carbohydrates. After two 20 minute washes in 3% acetic acid, the gels were visualized using a SYBR-green filter and an Alpha Innotech imaging system in movie mode. Following acquisition of the emerald-green signal, gels were de-stained overnight in 3% acetic acid and subsequently stained with SYPRO Ruby fluorescent protein stain (Invitrogen) and visualized using an ethidium bromide filter as above.

Antibody- and Complement-Binding Assays

To assess binding of Ab or complement to the surface of *F. tularensis*, 5×10<sup>8</sup> bacteria in 25 µl were mixed with 25 µl

of sterile, pre-cleared (10,000×g for 10 min) 0.5×PBS containing 0.1% glucose and either i) 1-5 µl (empirically-determined) of the test Ab/sera (sera were heat-inactivated) for Ab-binding assays or ii) 12.5 µl of intact or heat-inactivated normal mouse sera (for complement-binding assays). To assess the impact of protease inhibitors on Ab-binding, a protease inhibitor cocktail (Sigma, p2714—which includes aprotinin, a plasmin inhibitor) was included in the above Ab mixture. Samples were incubated at 37° C. for 1.5 hrs (for Ab-binding assays) or for various times (10 min-1 hr) for complement-binding assays. Following incubation, bacteria were pelleted by centrifugation (10,000×g for 10 min) and 45 µl of supernatant was removed. The bacteria were gently resuspended in 1 ml of 0.5×PBS containing 0.1% glucose and pelleted by centrifugation (10,000×g for 10 min) after which the supernatants were aspirated. Following one additional wash, the cells were resuspended to 25 µl in 50 mM Tris pH 8.0 containing protease inhibitors followed by the addition of 25 µl of sample buffer prior to boiling and SDS-PAGE. 1×10<sup>8</sup> bacteria per lane were resolved by SDS-PAGE and probed for bound immunoglobulin (IgG or IgM) heavy chain (HC) by western blot. Following development of the Ig HC signals, we re-probed the membranes for total FopA and quantified the data as surface Ab/total FopA and normalized the ratios to the corresponding MHB result. Statistical analysis was performed with the 2-tailed, T-test with Bonferroni corrections when appropriate and significance set at p<0.05. As a control, we included a BHI-grown wbtA strain [114] since such strains produce neither LPS OAg nor capsular OAg [30]. Normal mouse serum (NMS) and α-IgIC Ab (directed against a sub-surface protein) were used as control Abs.

#### Immunization and Challenge

C57Bl6 mice (Taconic, Germantown N.Y.) were maintained and bred in a specific pathogen free environment in the Animal Resource Facility at Albany Medical College. All experiments were conducted using 6-8 week old mice of both sexes and all the animal procedures conformed to the Institutional Animal Care and Use Committee guidelines.

Mice were immunized and boosted i.n. with 4×10<sup>7</sup> iFt LVS per mouse on days 0, 14 and 28. Test sera were drawn 12 days after each immunization and pooled; pooled sera was used in western blot analysis at a 1:500 dilution to probe BHI-grown Ft LVS.

On day 42, mice were anesthetized by intra-peritoneal injection of 100 µl of xylazine (20 mg/ml) and ketamine (1 mg/ml) and challenged by intranasal instillation of ~20 CFU of *F. tularensis* SchuS4 in 40 µl of PBS (equally divided between the two nostrils.)

Mice were monitored once-to-thrice daily for 21 days. No deaths occurred after day 8; PBS-immunized control mice succumb by day 7. Survival data are shown in the Kaplan Meier plot of FIG. 3B.

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## SEQUENCE LISTING

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&lt;223&gt; OTHER INFORMATION: synthetic linker peptide

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&lt;213&gt; ORGANISM: Artificial Sequence

&lt;220&gt; FEATURE:

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What is claimed is:

1. A nucleic acid that encodes a fusion protein, said nucleic acid comprising a first polynucleotide that consists of the nucleotide sequence of SEQ ID NO: 2; and a second polynucleotide sequence that consists of the nucleotide sequence of SEQ ID NO: 3.

2. A nucleic acid of claim 1 comprising the nucleotide sequence of SEQ ID NO: 1.

3. The nucleic acid of claim 1, wherein said first polynucleotide is linked to said second polynucleotide by a third polynucleotide that encodes a peptide linker.

4. The nucleic acid of claim 1, wherein said first polynucleotide is covalently linked to two or more second polynucleotides in tandem.

5. The nucleic acid of claim 3, wherein said peptide linker is selected from the group consisting of: a glycine linker, a glycine-rich linker and a glycine-serine linker.

6. The nucleic acid of claim 3, wherein the linker is about 3 to about 15 amino acids in length.

7. The nucleic acid of claim 3, wherein the linker is about 5 to about 10 amino acids in length.

8. A vector comprising the nucleic acid of claim 1.

9. An isolated cell comprising the nucleic acid of claim 1.

10. The isolated cell of claim 9, wherein said cell is a bacterial cell.

11. The isolated cell of claim 10, wherein said bacterial cell is selected from the group consisting of *Escherichia coli*, *Acinetobacter baumannii*, *Mycobacterium tuberculosis*, *Borrelia burgdorferi*, *Staphylococcus aureus*, *Yersinia pestis* and *F. tularensis*.

12. The isolated cell of claim 10, wherein said bacterial cell is selected from the group consisting of *Escherichia coli*, *Yersinia pestis*, and *F. tularensis*.

13. The isolated cell of claim 10, wherein said bacterial cell is *F. tularensis*.

14. The isolated cell of claim 9, wherein said isolated cell expresses a Tul4A-C3d(n) fusion protein, wherein n=1 to 5.

15. A vaccine comprising an isolated cell of claim 9 and a pharmaceutically acceptable carrier.

16. The vaccine of claim 15, further comprising an adjuvant.

17. A nucleic acid that encodes a fusion protein, the nucleic acid comprising:

(a) a groEL promoter; and

(b) the nucleic acid of claim 1.

18. A vector comprising the nucleic acid of claim 17.

19. An isolated bacterial cell comprising the nucleic acid of claim 17.

20. The isolated cell of claim 19, wherein said bacterial cell is selected from the group consisting of *Escherichia coli*, *Acinetobacter baumannii*, *Mycobacterium tuberculosis*, *Borrelia burgdorferi*, *Staphylococcus aureus*, *Yersinia pestis* and *F. tularensis*.

21. The isolated cell of claim 19, wherein said bacterial cell is selected from the group consisting of *Escherichia coli*, *Yersinia pestis* and *F. tularensis*.

22. The isolated bacterial cell of claim 19, wherein said cell is *F. tularensis*.

23. A vaccine comprising the bacterial cell of claim 19 and a pharmaceutically acceptable carrier.

24. A method of enhancing the immunogenicity of a bacterial cell, the method comprising: transforming said bacterial cell with the vector of claim 8 and growing the transformed bacterial cell under conditions favorable for expression of the fusion protein by the bacterial cell.

25. A method for immunizing a mammal against tularemia, the method comprising administering the vaccine of claim 15.

26. A method of enhancing the immunogenicity of a bacterial cell, the method comprising: transforming said bacterial cell with the vector of claim 18 and growing the transformed bacterial cell under conditions favorable for expression of the fusion protein by the bacterial cell.

27. A method for immunizing a mammal, the method comprising contacting said mammal with an isolated bacterial cell of claim 19.

\* \* \* \* \*